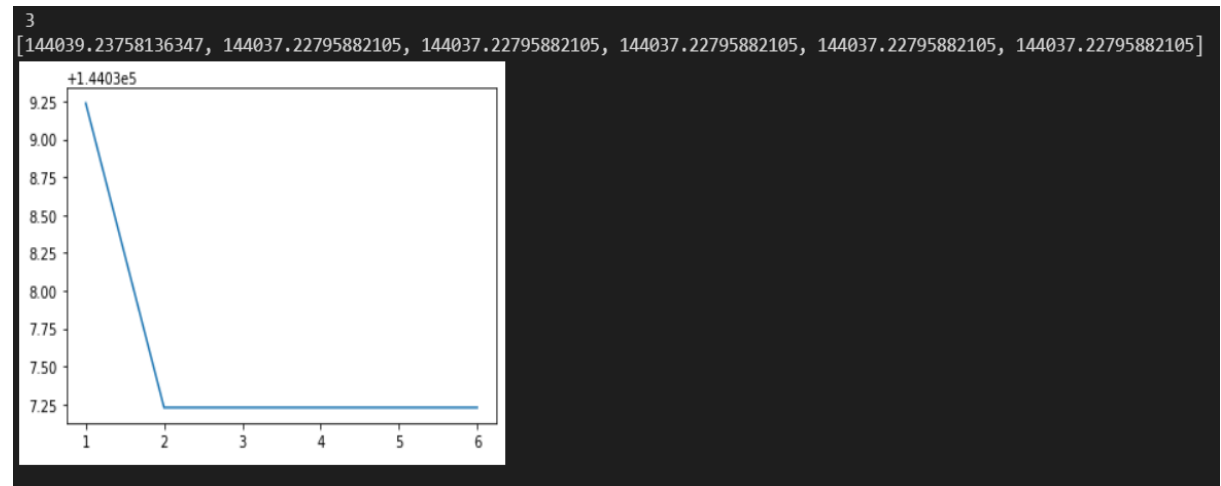


2. a. (i)

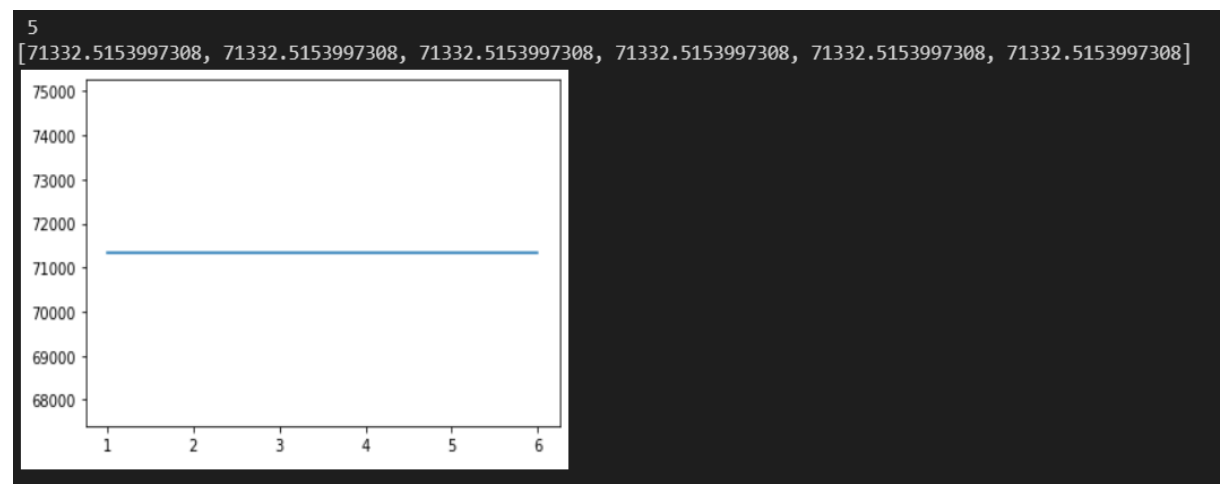
For $k = 3$

Total SSE of each clustering run



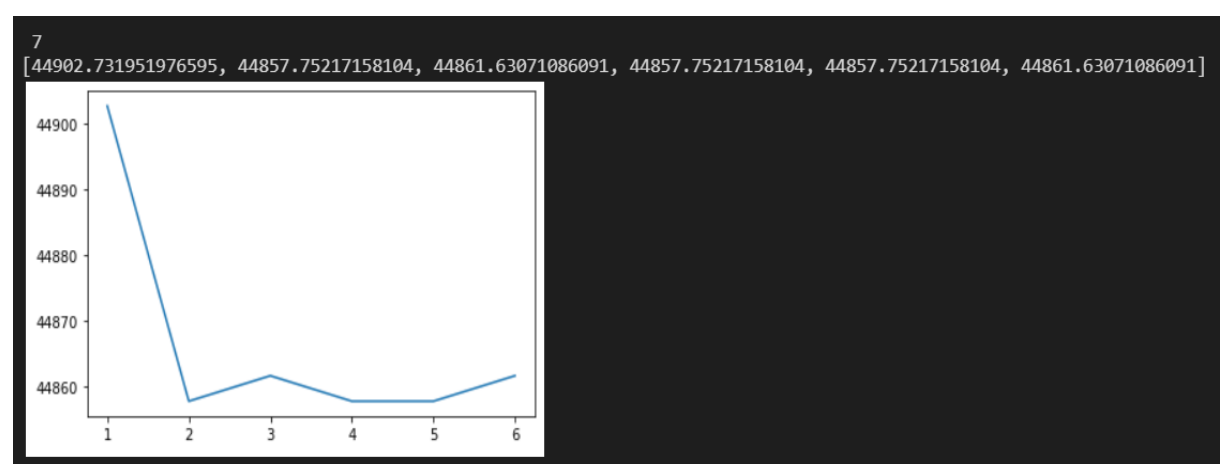
For $k = 5$

Total SSE of each clustering run



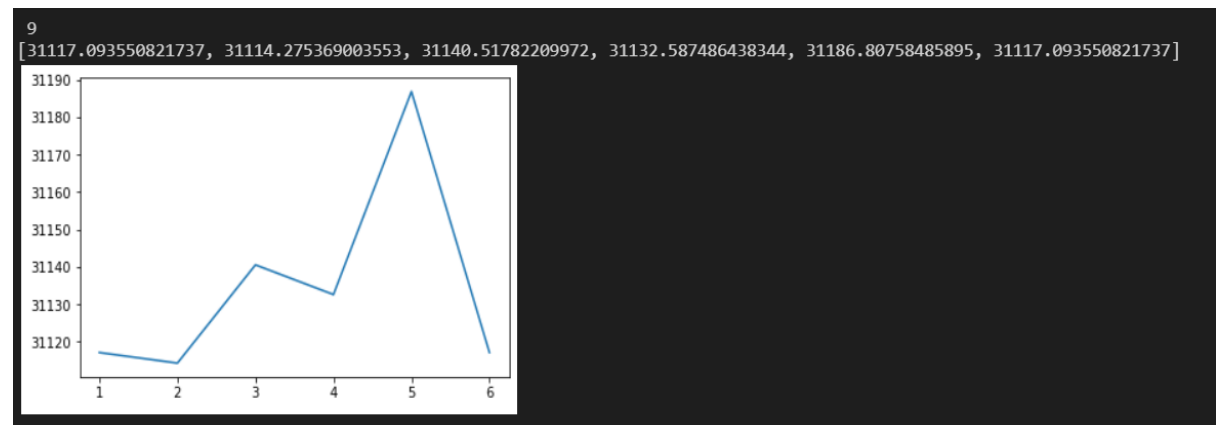
For $k = 7$

Total SSE of each clustering run



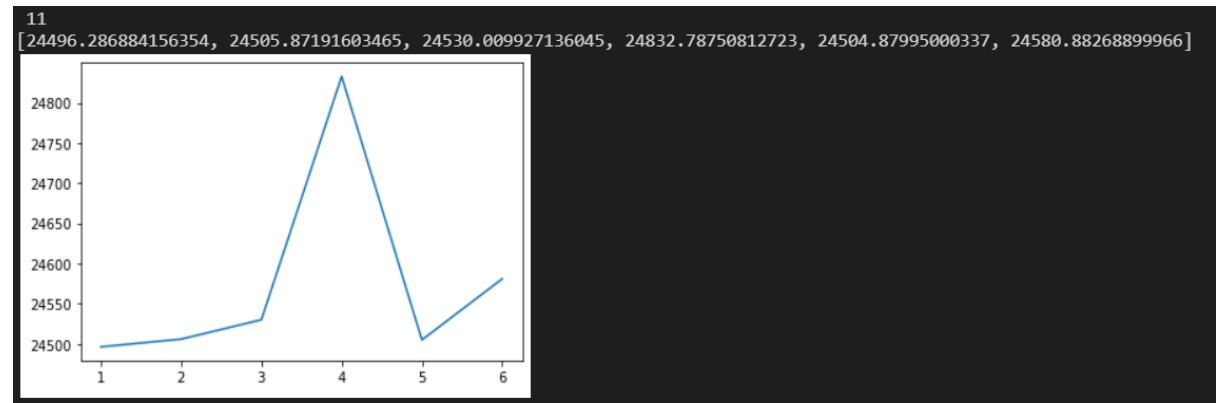
For $k = 9$

Total SSE of each clustering run

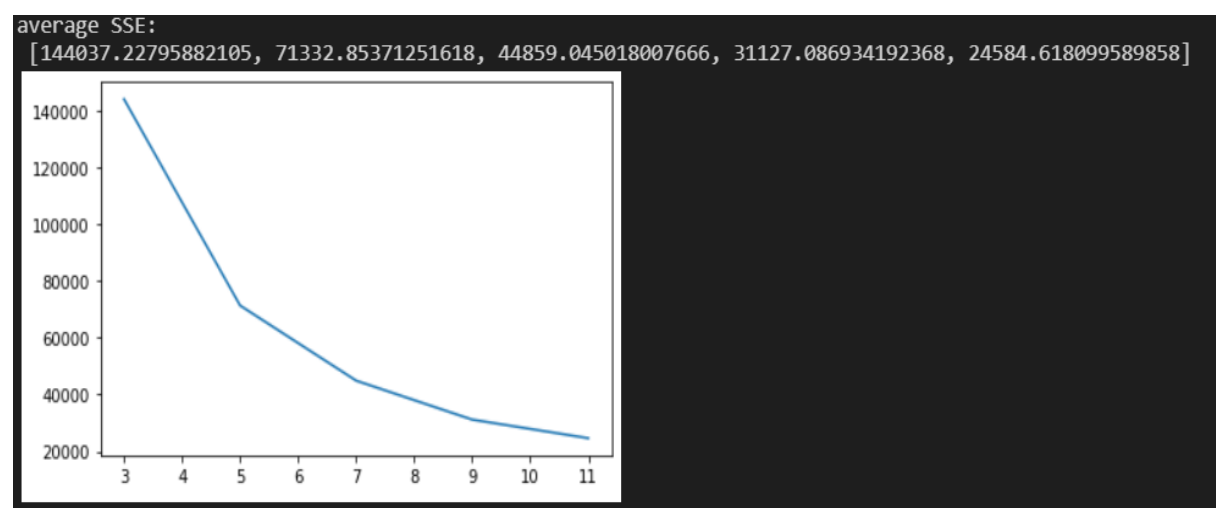


For $k = 11$

Total SSE of each clustering run



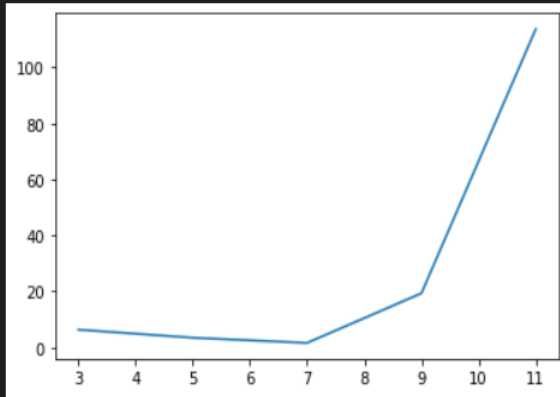
(ii) Average SSE for each k



Standard deviation of SSE for each k

std dev of SSE for each k:

[6.297677097707217, 3.4159507264437656, 1.5673954502974363, 19.312423449152938, 113.75987884365036]

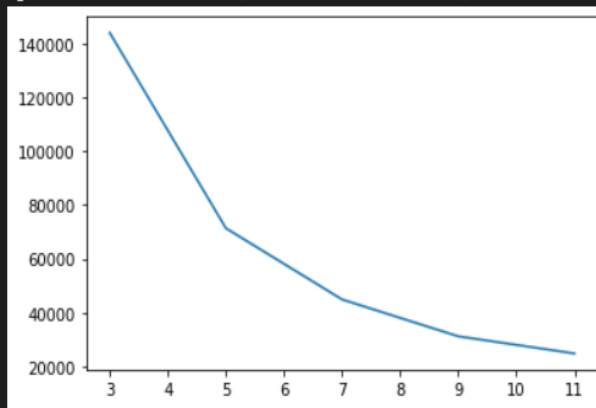


(iii) Min and max SSE for each k

Max SSE for each k

max SSE for each k:

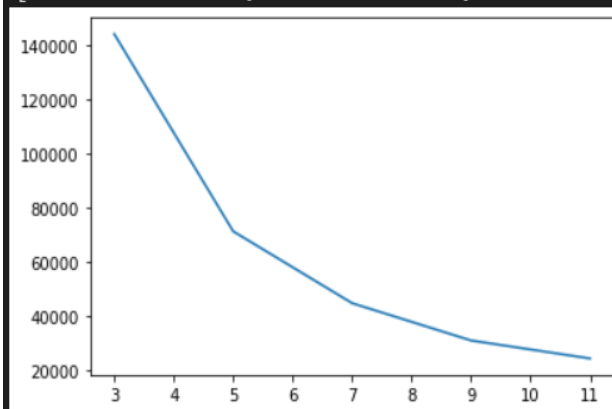
[144039.23758136347, 71341.04890438446, 44857.8551999683, 31156.266077488617, 24789.042556625303]



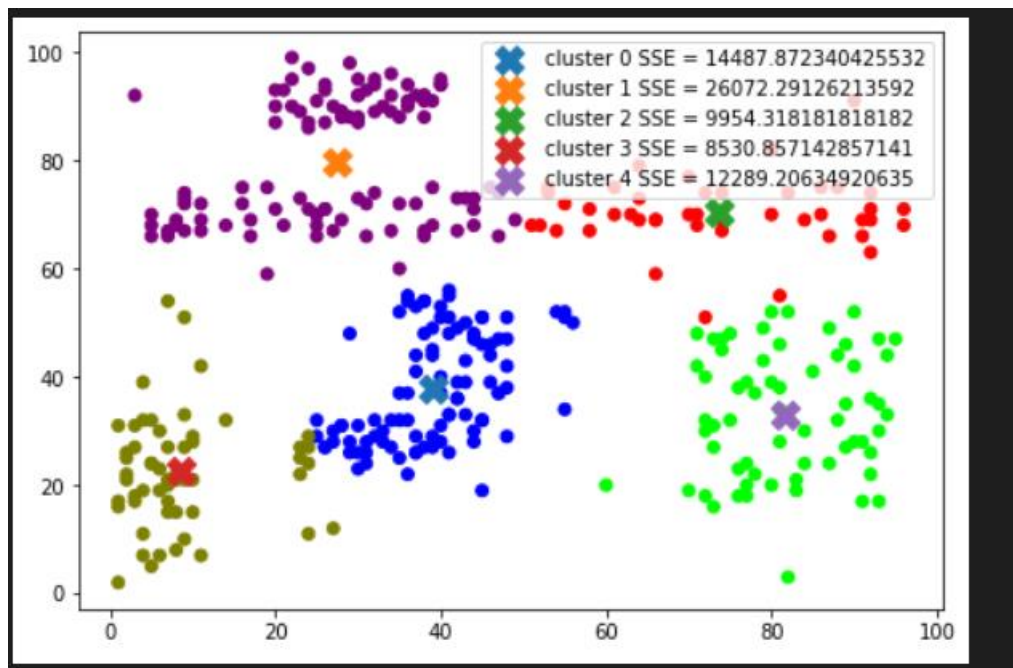
Min SSE for each k

min SSE for each k:

[144037.22795882105, 71332.5153997308, 44857.75217158104, 31114.275369003553, 24492.17944919884]



b.



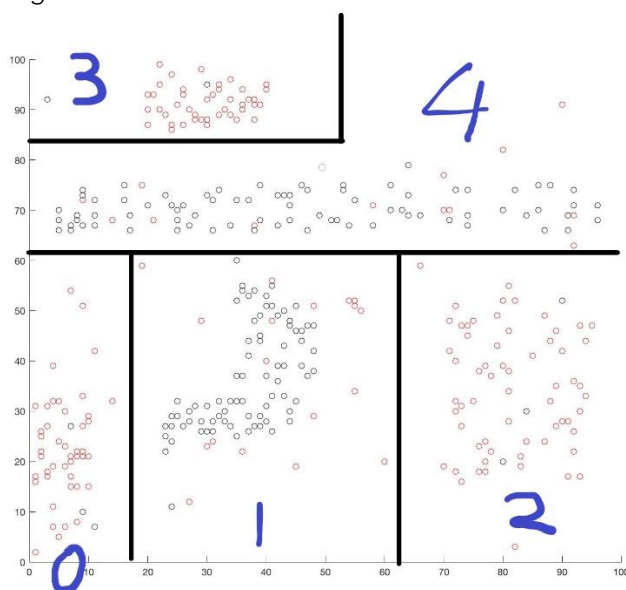
c.

The top cluster should be separated with the belt cluster, and the belt region in the middle should be in a single cluster.

Because the Euclidean distance is used, kmeans cannot correctly cluster the belt region in the middle. This proved that the k means clustering doesn't work well if the data is not spherically distributed.

d.

These data can be classified with linear boundaries, since the margin between each cluster is rather large.



```

man_class = []
for i in range(360):
    if df.iloc[i, 0] <= 18 and df.iloc[i, 1] <= 61:
        man_class.append(0)
    elif 18 < df.iloc[i, 0] <= 62 and df.iloc[i, 1] <= 61:
        man_class.append(1)
    elif df.iloc[i, 0] > 62 and df.iloc[i, 1] <= 61:
        man_class.append(2)
    elif df.iloc[i, 0] <= 58 and df.iloc[i, 1] > 84:
        man_class.append(3)
    else:
        man_class.append(4)
m_class = np.array(man_class)

#plt.scatter(df.loc[:, 0], df.loc[:, 1], c = man_class, cmap = "brg")
#plt.show()

```

e.

Construct a contingency matrix, which is symmetric

k_means = cluster

manual clustering = class

nij = intersection(class i, cluster j)

nij = nji

class/cluster	0	1	2	3	4
0	n00	n01	n02	n03	n04
1		n11	n12	n13	n14
2			n22	n23	n24
3				n33	n34
4					n44

$\text{rand_idx} = (a + d) / (a + b + c + d)$

$a = \text{sigma}(\text{comb}(n_{ij}, 2))$

$b = \text{sigma}(\text{comb}(n_{i.}, 2)) - \text{sigma}(\text{comb}(n_{ij}, 2))$

$c = \text{sigma}(\text{comb}(n_{.j}, 2)) - \text{sigma}(\text{comb}(n_{ij}, 2))$

$d = \text{comb}(N, 2) - a - b - c$

N is the total number of data points

rand index = $(a + d) / (a + b + c + d) = 0.8963478799133395$

The rand index represents the accuracy of the target clustering, which is kmeans in this case.

(Or the similarity between two clustering.)

```
a = 0
b = 0
c = 0
d = 0

for j in range(5):
    for i in range(5):
        m_ij = m_class[km.labels_ == j]
        nij = m_ij[m_ij == i].size
        if nij >= 2:
            a += spy.comb(nij, 2)

for i in range(5):
    nidot = m_class[m_class == i].size
    if nidot >= 2:
        b += spy.comb(nidot, 2)
b = b - a

for j in range(5):
    ndotj = km.labels_[km.labels_ == j].size
    if ndotj >= 2:
        c += spy.comb(ndotj, 2)
c = c - a

d = spy.comb(360, 2) - a - b - c

randidx = (a + d) / (a + b + c + d)
print(randidx)
```